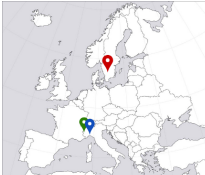


# Spatial Diffusion in SIR type Models: Simulation for Covid-19 Data

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# Introduction



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**2022**

Bachelor  
Degree in  
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**2023-2024**

Tutoring in  
Numerical  
Analysis and  
Calculus



**2024**

Master  
Degree in  
Mathematics  
at UniTO



**2024-2025**

Fellowship in  
Biostatistics at  
the Clinical  
Department  
Science - UniTO

# Introduction

## Skills

Bio-mathematics and Bio-statistics,  
Dynamical Systems, Complex Systems,  
Neural Networks, MonteCarlo  
Simulations, Assurance and Financial  
Mathematic

### Modeling



### Coding

R, Python, MATLAB, Maple, C++

LaTeX, Git, manim (python graphic  
tool).  
English language, level C1

### Programs and Languages



# The SIR model

**Total Population:**  $N = S(t) + I(t) + R(t)$

● Infected (I) ● Susceptibles (S) ● Removed (R)

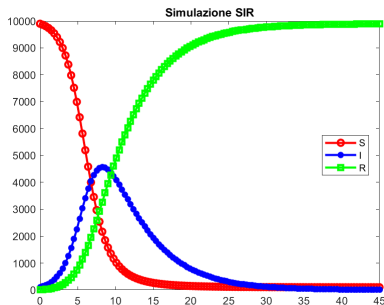


Figure: Numeric Simulation of the SIR model

## Model Equations

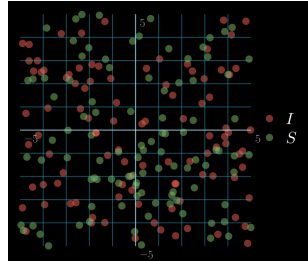
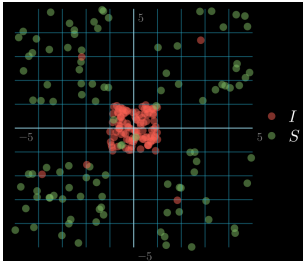
$$\begin{cases} \dot{S} = -\frac{\beta}{N}S(t)I(t) \\ \dot{I} = -\frac{\beta}{N}S(t)I(t) - \gamma I(t) \\ \dot{R} = \gamma I(t) \end{cases}$$

- ✓  $\beta$  is the infection rate  $I$
- ✓  $\gamma \propto T^{-1}$  where  $T$  is the average time of recovery from the illness.

# An Hidden Assumption

## Perfect Mixing

$S$  and  $I$  are uniformly distributed in space at all times

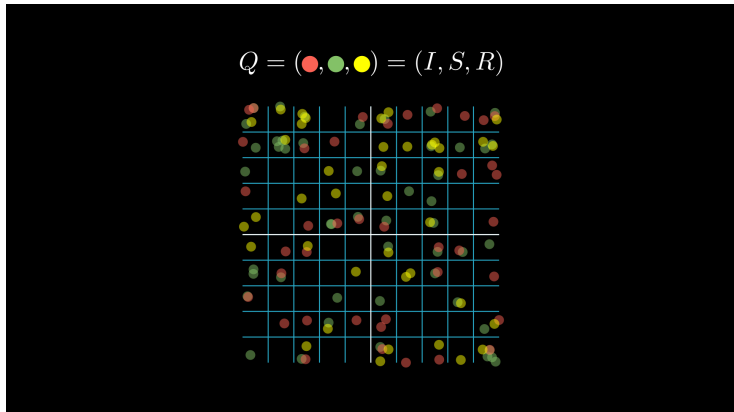


**Figure:** In the initial phases of an epidemic diffusion, the infected population  $I$  is not uniformly spread across space

# Lattice Gas Cellular Automata (LGCA)

## Particles and Cells

Particles moving on a lattice can interact only inside the same cell [Schneckenreither et al.]



# Particles Laws of Motion

At each time step  $n$  particles move to a neighbor cell

## Uniqueness

At most one particle can reach a fixed neighbor cell

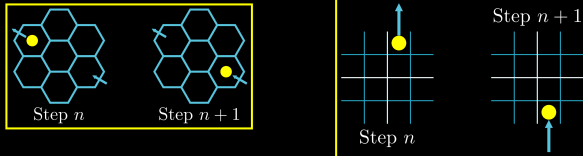
## Randomness

The exit configuration is randomly selected

## Motion's Directions



## PacMan Effect



# Epidemic Laws described

## Evolution Rules

- Each particle of state  $I$  of a specific cell, infects a particle of state  $S$  of the same cell with probability  $\beta_{LG}$
- With rate  $\gamma$  particles pass from state  $I$  to state  $R$

## Epidemic Laws inside a cell

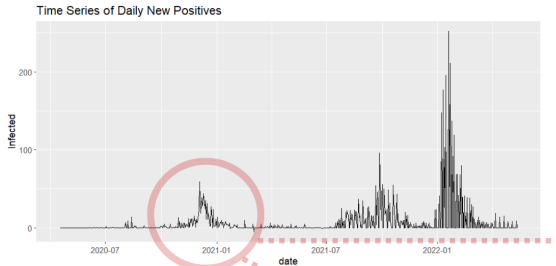
Let  $I(c)$  be the number of infected in cell  $c$  and let  $q_n(p)$  be the state of particle  $p \in c$  at time step  $n$ , then

$$\begin{cases} \phi_c = \mathbb{P}(q_{n+1}(p) = S | q_n(p) = S) = (1 - \beta_{LG})^{I(c)} \\ \psi_c = \mathbb{P}(q_{n+1}(p) = I | q_n(p) = S) = 1 - \phi_c \\ \mathbb{P}(q_{n+1}(p) = I | q_n(p) = I) = 1 - \beta_{LG} \\ \mathbb{P}(q_{n+1}(p) = R | q_n(p) = I) = \beta_{LG} \end{cases}$$

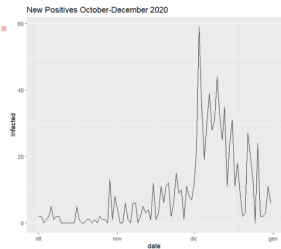


# Covid-19 in Kodiak Island, Alaska

## An overlook of data and estimated parameters

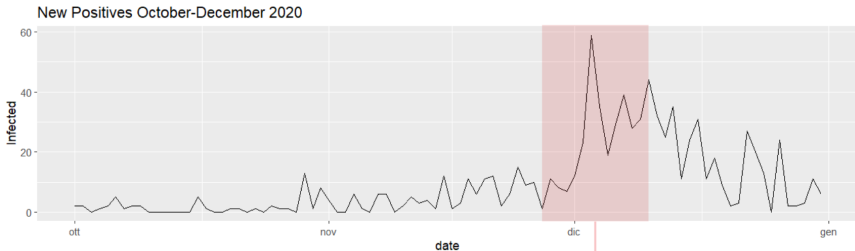


Parameter	Value
<chr>	<chr>
Starting Date	2020-10-01
Total Population	13100
Initial Population of Infected People	15
gamma	0.153846153846154
beta	0.34588132363107



# Covid-19 in Kodiak Island, Alaska

## How to estimate $\beta$ from new confirmed cases

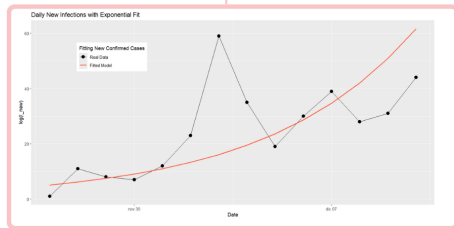


Assuming  $S \approx N$  in the beginning of epidemic peak

$$\frac{dI}{dt} = \frac{\beta}{N} SI - \gamma I \approx (\beta - \gamma)I$$

Hence, if  $C$  is the number of new infected each day, we have

$$C = C_0 e^{(\beta - \gamma)t}$$



# Covid-19 in Kodiak Island, Alaska

## SIR and LGCA Simulation Results

### LGCA

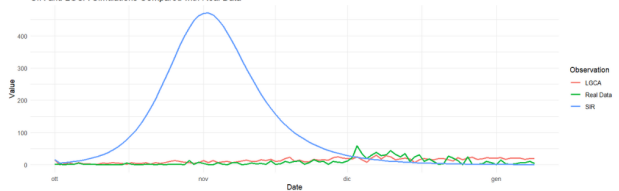
Parameter <chr>	Value <chr>
Starting Date	2020-10-01
Population	12996
L <sub>0</sub>	15
gamma	0.153846153846154
beta	0.34588132363107
cells	12996

### SIR

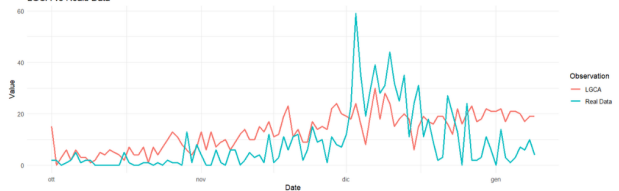
Parameter <chr>	Value <chr>
Starting Date	2020-10-01
Population	13100
L <sub>0</sub>	15
gamma	0.153846153846154
beta	3.23807026513713e-05

$$\beta_{LG} = \beta_{SIR} \cdot cells$$

SIR and LGCA Simulations Compared with Real Data

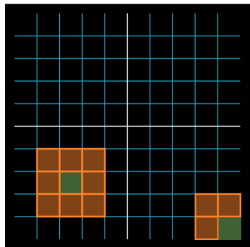


LGCA vs Reald Data

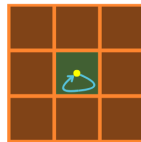
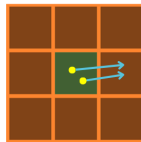
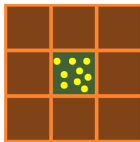


# Modified LGCA algorithm

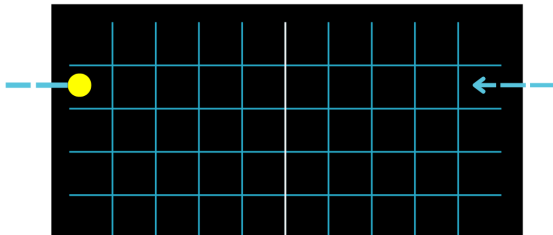
Neighbors



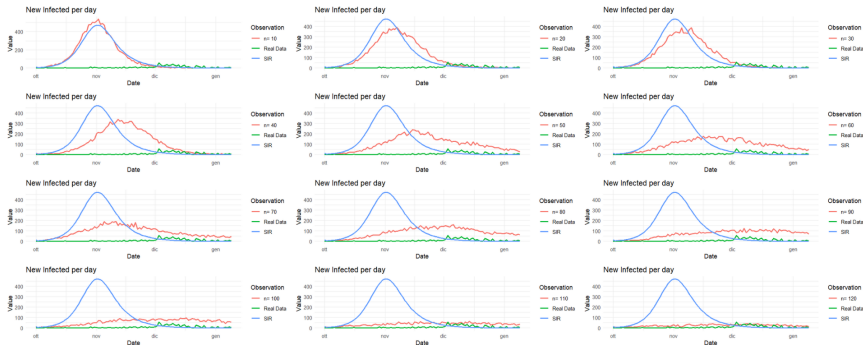
Yes



No



# Modified LGCA Simulations

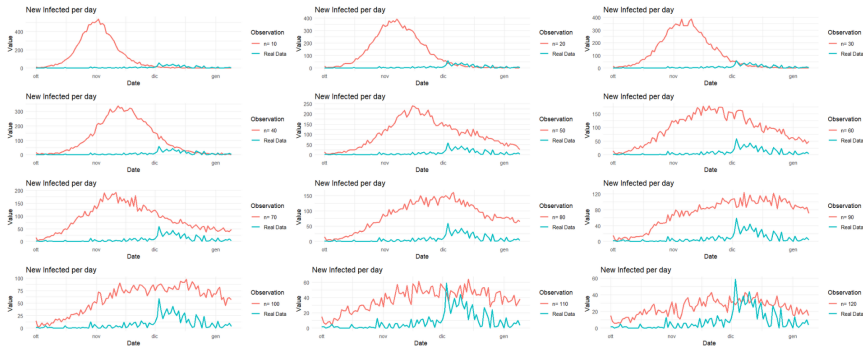


cells	beta_LG
100	0.002640315
400	0.010561262
900	0.023762839
1600	0.042245047
2500	0.066007886
3600	0.095051356
4900	0.129375457
6400	0.168980189
8100	0.213865551
10000	0.264031545

Parameter	Value
<chr>	<chr>
Starting Date	2020-10-01
Population	13100
L_0	15
gamma	0.153846153846154
beta	3.23807026513713e-05

$$\beta_{LG} = \beta_{SIR} \cdot \text{cells}$$

# Modified LGCA Simulations

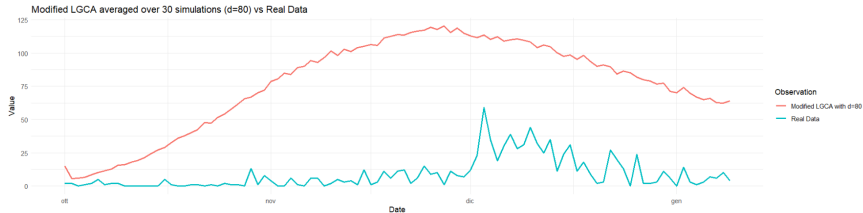


cells	beta_LG
100	0.002640315
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Parameter	Value
<chr>	<chr>
Starting Date	2020-10-01
Population	13100
L_0	15
gamma	0.153846153846154
beta	3.23807026513713e-05

$$\beta_{LG} = \beta_{SIR} \cdot cells$$

# Modified LGCA Simulations



Parameter <chr>	Value <chr>
Starting Date	2020-10-01
Number of Simulations	30
Lattice Dimension	6400
beta_LG	0.168980188644187
gamma	0.153846153846154
I_0	15
Population	13100

# Final Observations

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## Issues

Parameters and Initial Values Estimation,  
Dataset Choice, Small Number of Simulations.

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## Open Questions

Link between parameters (e.g. lattice dimension) and qualitative behaviour of Modified LGCA, Stability of the solutions and dependence on distributions and laws of motion

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## Conclusions

Modified LGCA seems to provide promising results into using lattice based structures to model spatial diffusion in epidemic peaks, specially for small populated areas

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# Bibliography



Günter Schneckenreither, Nikolas Popper, Günther Zauner, and Felix Breitenecker.

Modelling sir-type epidemics by odes, pdes, difference equations and cellular automata—a comparative study.

*Simulation Modelling Practice and Theory*, 16(8):1014–1023, 2008.



Junling Ma.

Estimating epidemic exponential growth rate and basic reproduction number.

*Infectious Disease Modelling*, 5:129–141, 2020.